|                           | F Er. S Corrected by the ST  |                                 |                     |
|---------------------------|--|---------------------------------|---------------------|
| umb r: 09/896             | 0,779  | CRF Processing Da Edited by:    | 1                   |
| Changed a file from no    | on-ASCII to ASCII  | Verified by:                    | (STIC sta           |
| Changed the margins i     | n cases where the sequence text wa   | as "wrapped" down to the next   | line.               |
| Edited a format error in  | the Current Application Data section                                       | n, specifically:                |                     |
|                           | lication Data section with the actual oprior application data; or 🔲 other  | current number. The number in   | nputted by the      |
| Added the mandatory h     | neading and subheadings for "Curren  | nt Application Data".           |                     |
| Edited the "Number of     | Sequences" field. The applicant spel                                       | lled out a number instead of us | sing an integer.    |
| Changed the spelling o    | f a mandatory field (the headings or s                                     | subheadings), specifically:     |                     |
| Corrected the SEQ ID N    | NO when obviously incorrect. The se  | equence numbers that were ed    | ited were:          |
| Inserted or corrected a   | nucleic number at the end of a nuclei                                      | ic line. SEQ ID NO's edited:    |                     |
|                           | placement. All responses must be on<br>onse below the subheading, this was |                                 |                     |
| Inserted ∞lons after he   | eadings/subheadings. Headings edit   | ed included: ,                  |                     |
| Deleted extra, invalid, h | neadings used by an applicant, specif                                      | fically:                        |                     |
|                           | I "garbage" at the beginning/end of file bughout text;                     |                                 | ame at end of file; |
| Inserted mandatory hea    | adings, specifically:  |                                 |                     |
| Corrected an obvious e    | error in the response, specifically:                                       |                                 |                     |
| Edited identifiers where  | upper case is used but lower case is                                       | s required, or vice versa.      | •                   |
| Corrected an error in th  | e Number of Sequences field, specif  | ically:                         |                     |
| A "Hard Page Break" co    | ode was inserted by the applicant. A                                       | Il occurrences had to be delete | ed.                 |
|                           | don in amino acid sequences and ad<br>Sequences corrected:                 |                                 |                     |
| Other:                    | •  |                                 |                     |
| •                         | <del></del>  |                                 |                     |
|                           |  |                                 |                     |
| _                         |  |                                 |                     |

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95



PCT09

DATE: 04/30/2002 RAW SEQUENCE LISTING PATENT APPLICATION: US/09/890,779 TIME: 15:48:33

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\04302002\1890779.raw

4 <110> APPLICANT: WOLFGANG WERR 6 <120> TITLE OF INVENTION: A METHOD FOR INHIBITING THE EXPRESSION OF TARGET GENES IN PLANTS 10 <130> FILE REFERENCE: A34537-PCT-USA 12 <140> CURRENT APPLICATION NUMBER: 09/890,779 13 <141> CURRENT FILING DATE: 2001-08-06 15 <150> PRIOR APPLICATION NUMBER: PCT/EPOO/01524 16 <151> PRIOR FILING DATE: 2000-02-09 18 <150> PRIOR APPLICATION NUMBER: EP99420030.1 19 <151> PRIOR FILING DATE: 1999-02-09 21 <160> NUMBER OF SEQ ID NOS: 2 23 <170> SOFTWARE: FastSEQ for Windows Version 4.0 25 <210> SEQ ID NO: 1 26 <211> LENGTH: 6 27 <212> TYPE: DNA 28 <213> ORGANISM: BACTERIA 30 <400> SEQUENCE: 1 6 31 gaattc 33 <210> SEQ ID NO: 2 34 <211> LENGTH: 8 35 <212> TYPE: DNA 36 <213> ORGANISM: BACTERIA 38 <400> SEQUENCE: 2 8

39 gatctcga

VERIFICATION SUMMARYDATE: 04/30/2002PATENT APPLICATION: US/09/890,779TIME: 15:48:34

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\04302002\1890779.raw



## Does Not Comply Corrected Diskette Needed

PCT09

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/890,779

DATE: 04/25/2002 TIME: 12:01:12

Input Set : A:\34537 sequence listing.txt
Output Set: N:\CRF3\04252002\1890779.raw

4 <110> APPLICANT: WOLFGANG WERR

6 <120> TITLE OF INVENTION: A METHOD FOR INHIBITING THE EXPRESSION

7 OF TARGET GENES IN PLANTS

10 <130> FILE REFERENCE: A34537-PCT-USA

12 <140> CURRENT APPLICATION NUMBER: 09/890,779

13 <141> CURRENT FILING DATE: 2001-08-06

15 <150> PRIOR APPLICATION NUMBER: PCT/EPOO/01524

16 <151> PRIOR FILING DATE: 2000-02-09

18 <150> PRIOR APPLICATION NUMBER: EP99420030.1

19 <151> PRIOR FILING DATE: 1999-02-09

21 <160> NUMBER OF SEQ ID NOS: 2

23 <170> SOFTWARE: FastSEQ for Windows Version 4.0

## ERRORED SEQUENCES

33 <210> SEQ ID NO: 2

34 <211> LENGTH: 8

35 <212> TYPE: DNA

36 <213> ORGANISM: BACTERIA

38 <400> SEQUENCE: 2

39 gatctcga

E--> 46 (ny02:355030.1) - delete

VARIABLE LOCATION SUMMARY

DATE: 04/25/2002

PATENT APPLICATION: US/09/890,779

TIME: 12:01:13

Input Set : A:\34537 sequence listing.txt
Output Set: N:\CRF3\04252002\I890779.raw

## Use of n's or Xaa's(NEW RULES):

Use of n's and/or Xaa's have been detected in the Sequence Listing.
Use of <220> to <223> is MANDATORY if n's or Xaa's are present.
in <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

Seq#:2; N Pos. 9

## VERIFICATION SUMMARY

DATE: 04/25/2002 TIME: 12:01:13

PATENT APPLICATION: US/09/890,779

Input Set : A:\34537 sequence listing.txt Output Set: N:\CRF3\04252002\1890779.raw

L:46 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:2

L:46 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:12 SEQ:2

L:46 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:10

L:46 M:112 C: (48) String data converted to lower case,

L:46 M:252 E: No. of Seq. differs, <211> LENGTH:Input:8 Found:12 SEQ:2